The genetic code of Protein synthesis"

= 1st lecture "The genetic code"

\*\*Metabolism > Catabolism . "Cutting [3]"

Anabolism : Is to use the small nutrients to build the bulk units.

Anabolism of Carbohy drates can be done "easily" by reversing the catabolism 9 "pyruvic acid > glucose > glycogen"

\*\*How to build up proteins? 90?

This process happens after the catabolism "Digestion > absorption" inside your body "and gives amino acids, the the Cell decides what type of protein it needs" enzymes obuilding proteins.

• hormones • Secretions ...etc"

All that happens by the "Protein synthesis"

- In order to make proteins we need some reference..

  the reference is "DNA" Unucleic acid" which's found inside
  the nucleus of contains "nucleotides"
- Nucleotides carrangement is very precise, this arrangement determines the type of amino acids that produced.
- To make protein a certain gene in the DNA will become active.
- DNA inside the nucleus is a chromotin material "befor division".
   DNA become as chromosomes "During division".

46 chromosom => each contains one DNA molecule.



Coiling the DNA & shortining its shape is done by "Histories" proteins

Share & Care Group

Life = is protein & DNA contain the protein, so the reference of protein synthesis should be kept away from any-thing can affect it, this place is the "nucleus".

"expression: "the process of "transcription" to make protein.

Inside the chromosom the DNA is coiled but someplaces are linear, another places are more coiled and can't be "expressed"

\* Why is the coiling of DNA is shown more is some places than the others?"

A: Because these places are more connected to the "Exons"."

these highly coiled places can't be expressed.

Highly coiled places "not expressed" > Linear places "ppened" "Can be expressed" 3% only.

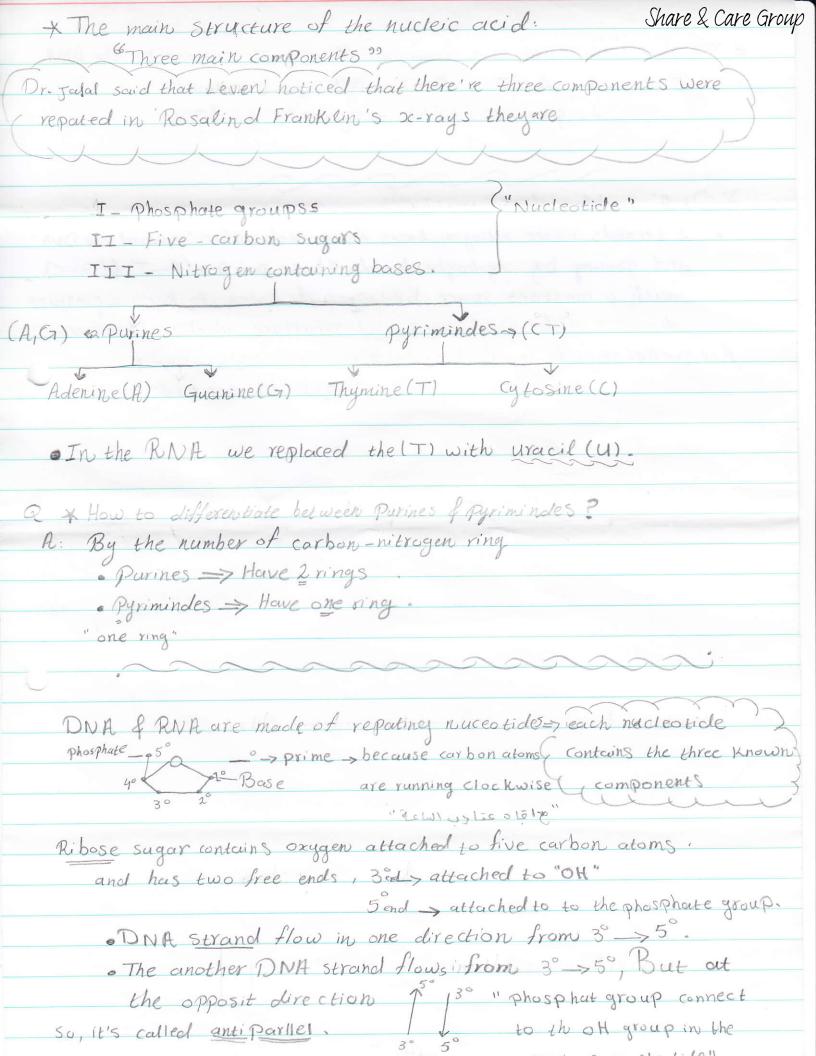
" sla Medit F"

\* Genetics showed before the DNA appearence.

important into sperm is only about => nucleus

· Ovule is a cell with a cytoplasm.

1221143



The four nucleotides are found in different proportions In the DNA But :- 1- A = T,  $G = C \rightarrow Allways$  /=:equals

2-purines (A&G) = Pyrimidnes (C&T).

## \* DNA double helix Structure

and pairing by hydrogen bonds (base pair) -> (A-T) f (G-C) with a constant space between the bases, to keep a constant diameter of the DNA antiparllel structure which's "2 nanometer"

Antiparllel strands in Arabic: show his base / chip for it has a constant with the bases of the DNA antiparllel structure which's "2 nanometer".

Antiparllel strands in Arabic: show his base / chip for it has a constant which is "to be the bases of the DNA antiparllel structure which's "2 nanometer".

The gene: Any part of the DWA can be expressed, bund in the places that are easy to open. (3%) or (7%)

\* Not all the DNA is genes, because we have highly coiled places, and as we said they can't be opened.

Inside your DNA 4 billion bases are found

\* However Knowing genes, and there places wasn't enough, so scientists determined the beginnings and ends of these gene \$00 because without Knowing them this science becomes useless or not understood. \*

By specifying where genes start dend, scientists could detect (find) the genetic diseases.

DNA is the only molecule that can replicat "copy" it self.
"In cell division"

\* How to differntiate between "Replication" of "Transcription"?

- Replication: Is making another DNA molecule for the daughter cells.

- Transcription: Is the synthesis of RNA from DNA template.

\* Women without her man is nothing = without the punctuation women without her, man is nothing => with the punctuation has an extreme different meaning.

Replication occurs inside the nucleus, the DNA is replicated but not the chromosome, means one chromosome can contain 2 DNA instead of one by replication the parent DNA to give the daughter strands.

- [Rule: The chromosome can't be replicated]-

The highly wild places which open only during the replication and open after the "replication origin", these highly could place are opened by enzymes.

"Replication origin": The linear places, and they open first,
which is a specific sequence of nucleotides.

By opening the replication origin, we have this shape

The DNA step by step starts to be opened and the two strands separate at the end, by specific proteins of enzymes.

| X Hydrogen bonds are weak so they can be be broken down by increasing the temprature or adding acid (changing PH).

2- After separating the parent strands, the nucleoticles that are already existed in the nucleus "come from catabolism of the digested food", they start to make "base pairing" with the daughter strands.

\* Replication of DNA in 5 steps

Share & Care Group

from each other

1. The Un winding proteins keep the two strands separated and away 1, the "Helicase" aspecific enzyme moves between the two strands and cut the hydrogen bonds.

"The result: two separated strands, which are kept separated by the unwinding proteins " these proteins remove the histories"

2 - Now in the "Building a primer", the nucleotides in nucleus try to start base pairing with the separated strands, because their nucleotides are able to make base pairing with the another nucleotides, to build up a "primer", which helps in DNA replication.

The primer. Is a sequence of (10-12) nucleotides, it's the basic unit in building the new DNA-

The requirments for building up a new DNA: 1- A primer: RNA "A short stretch of RNA". 2- RNA polymerase.

\* RNA polymerase bind the RNA and fixes it to the DNA strand, to make the building unit of the new DNA.

The two separated strands are:

1. Leading strand: 3° > 5°, its competated strand 5° -> 3°
2. Lagging strand: "Okazaki fragements" 5° -> 3°, its competated

Strand 3° -> 5°

1

\* The process of building rleading strand is easier than the lagging strand.

3 - Assembling complementry strands.

Building the new strands, the replication starts from 5->3° so it's reasy to form a continous copy, by using an RNA primer. But the okazaki fragments complemntally strands forming is harder or more difficult because another RNA primer is needed everytime that the process stops.

DNA polymeras III (3), that cutalyzes the formation of the complementary strand on the template strand, more specificily it helps the nucleotides to bind to the template strands to form the complementary strands.

4- Removing the primers

DNA polymerase! (1) remove the primers from the lagging strands, gaps now formed between the nucleotides, which are filled by the DNA polymerase 11 (1) that fills them with nucleotides.

5- Joining the okazaki fragments:

The DNA ligase joins " was" the fragments to the lagging strand and makes it as a continuous strand.

"Enzymes work together at the same time"

- The whole process occurs in the whole DNA.

- -"Semiconservative replication": The parellel molecule of DIVA gives two daughter molecule contains one strand from the parent, in each new chromosomes has one strand of the parent of the other is a complemntary strand, this way keeps the genetic material to
- DNA polymeras 1(1) doesn't remove the first RNA primer, which causes alot of faulty that result as genetic diseases like disorders in some enzymes or some of them agen't existed